

# CRF Errors Corrected by the STIC Systems Branch

PCT09

Serial Number: 09/087, 962

CRF Processing Date: \_\_\_\_\_  
 Edited by: \_\_\_\_\_  
 Verified by: \_\_\_\_\_ (STIC staff)

**ENTERED**

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning end of files: ☐ secretary initials/filename at end of file;  
☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted **ending** stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.

PCT09

## RAW SEQUENCE LISTING

DATE: 04/12/2001

PATENT APPLICATION: US/09/787,962

TIME: 10:50:30

Input Set : A:\Cpg.pto

Output Set: N:\CRF3\04122001\I787962.raw

3 <110> APPLICANT: The University of British Columbia et al.  
 5 <120> TITLE OF INVENTION: A PLANT LONG CHAIN FATTY ACID BIOSYNTHETIC ENZYME  
 7 <130> FILE REFERENCE: 4810-58563  
 C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/787,962  
 10 <141> CURRENT FILING DATE: 2001-03-22  
 12 <150> PRIOR APPLICATION NUMBER: PCT/CA00/00860  
 13 <151> PRIOR FILING DATE: 2000-07-22  
 15 <150> PRIOR APPLICATION NUMBER: US60/145,013  
 16 <151> PRIOR FILING DATE: 1999-07-22  
 18 <160> NUMBER OF SEQ ID NOS: 5  
 20 <170> SOFTWARE: PatentIn Ver. 2.0  
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 23 <211> LENGTH: 2509  
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 25 <213> ORGANISM: Arabidopsis sp.  
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 30 agagaaacgt taggagcaaa caaacaagaag agaaaagacg atgatgactg atgagagctt 180  
 31 taacaaaaaa ataaaatgag agagctcaac gggtagaatt gtgagacttg agagagtgtt 240  
 32 tcctatttaa ggcattgcat tagtggttat tacgagaatg ccaccgaacg agtacatatt 300  
 33 aatgtatagt atgttaatga tagtctaact aaaatttggg ttttattgaa atagaatttt 360  
 34 gtaagaataa tgaggatctg taatatagct ggatttgcat taaatcgtae gccgttggtt 420  
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 37 aatagtcaaa cttaggtttac atcataatgg gccaatcttt tgagttgtga ttgatctcca 600  
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 58 gtatttttgt aacaagaaag caatgttgat ccctaattgc ttgttttagg ttggtgggtc 1860

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Output Set: N:\CRF3\04122001\I787962.raw

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VERIFICATION SUMMARY

PATENT APPLICATION: US/09/787,962

DATE: 04/12/2001

TIME: 10:50:31

Input Set : A:\Cpg.pto

Output Set: N:\CRF3\04122001\I787962.raw

L:9 M:270 C: Current Application Number differs, Replaced Current Application Number